

WHAT IS CLAIMED IS:

1. An isolated polynucleotide comprising a member selected from the group consisting of:
 - (a) a polynucleotide having at least 80% sequence identity, as determined by the GAP algorithm under default parameters, to a polynucleotide selected from the group consisting of SEQ ID NOS: 1, 3, and 5;
 - (b) a polynucleotide encoding a polypeptide selected from the group consisting of SEQ ID NOS: 2, 4, and 6;
 - (c) a polynucleotide amplified from a *Zea mays* nucleic acid library using primers which selectively hybridize, under stringent hybridization conditions, to loci within a polynucleotide selected from the group consisting of SEQ ID NOS: 1, 3, and 5;
 - (d) a polynucleotide which selectively hybridizes, under stringent hybridization conditions and a wash in 0.1X SSC at 60°C, to a polynucleotide selected from the group consisting of SEQ ID NOS: 1, 3, and 5;
 - (e) a polynucleotide selected from the group consisting of SEQ ID NOS: 1, 3, and 5;
 - (f) a polynucleotide which is complementary to a polynucleotide of (a), (b), (c), (d), or (e); and
 - (g) a polynucleotide comprising at least 30 contiguous nucleotides from a polynucleotide of (a), (b), (c), (d), (e), or (f).

Sub 3 2. A recombinant expression cassette comprising a member of claim 1 operably linked, in sense or anti-sense orientation, to a promoter.

3. A host cell comprising the recombinant expression cassette of claim 2.

Sub 3 4. A transgenic plant comprising a recombinant expression cassette of claim 2.

5. The transgenic plant of claim 4, wherein said plant is a monocot.

6. The transgenic plant of claim 4, wherein said plant is a dicot.

Sub A57
7. The transgenic plant of claim 4, wherein said plant is selected from the group consisting of: maize, soybean, sunflower, sorghum, canola, wheat, alfalfa, cotton, rice, barley, and millet.

Sub C47
8. A transgenic seed from the transgenic plant of claim 4.

9. A method of modulating the level of maize RAD51 in a plant, comprising:

- Sub A67*
- (a) introducing into a plant cell a recombinant expression cassette comprising a maize RAD51 polynucleotide of claim 1 operably linked to a promoter;
 - (b) culturing the plant cell under plant cell growing conditions;
 - (c) regenerating a whole plant which possesses the transformed genotype; and
 - (d) inducing expression of said polynucleotide for a time sufficient to modulate the level of maize RAD51 in said plant.

10. The method of claim 9, wherein the plant is maize.

11. An isolated protein comprising a member selected from the group consisting of:

- (a) a polypeptide of at least 20 contiguous amino acids from a polypeptide selected from the group consisting of SEQ ID NOS: 2 and 4;
 - (b) a polypeptide selected from the group consisting of SEQ ID NOS: 2 and 4;
 - (c) a polypeptide having at least 80% sequence identity to, and having at least one linear epitope in common with, a polypeptide selected from the group consisting of SEQ ID NOS: 2 and 4, wherein said sequence identity is determined using the GAP program under default parameters; and
 - (d) at least one polypeptide encoded by a member of claim 1.
- AA 27*